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RAW SEQUENCE LISTING
PATENT APPLICATION *US/09/175,713*

DATE: 11/03/98
TIME: 15:35:12

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This Raw Listing contains the General
Information Section and up to the first 5 pages.

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1                              SEQUENCE LISTING
2
3    (1)    General Information:                                    ENTERED
4
5         (i) APPLICANT: Herrmann, Stephen H.
6                       Lu, Zhijian
7                       McCoy, John M.
8                       Swanberg, Stephen L.
9                       Walker, Bruce
10                      Yang, Otto
11
12       (ii) TITLE OF INVENTION: CHEMOKINES WITH AMINO-TERMINAL MODIFICATIONS
13
14      (iii) NUMBER OF SEQUENCES: 15
15
16       (iv) CORRESPONDENCE ADDRESS:
17            (A) ADDRESSEE: Genetics Institute, Inc.
18            (B) STREET: 87 CambridgePark Drive
19            (C) CITY: Cambridge
20            (D) STATE: MA
21            (E) COUNTRY: U.S.A.
22            (F) ZIP: 02140
23
24        (v) COMPUTER READABLE FORM:
25            (A) MEDIUM TYPE: Floppy disk
26            (B) COMPUTER: IBM PC compatible
27            (C) OPERATING SYSTEM: PC-DOS/MS-DOS
28            (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
29
30       (vi) CURRENT APPLICATION DATA:
31            (A) APPLICATION NUMBER:
32            (B) FILING DATE:
33            (C) CLASSIFICATION:
34
35     (viii) ATTORNEY/AGENT INFORMATION:
36            (A) NAME: Sprunger, Suzanne A.
37            (B) REGISTRATION NUMBER: 41,323
38
39       (ix) TELECOMMUNICATION INFORMATION:
40            (A) TELEPHONE: (617) 498-8284
41            (B) TELEFAX: (617) 876-5851
42
43
44   (2) INFORMATION FOR SEQ ID NO:1:
45
46       (i) SEQUENCE CHARACTERISTICS:
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RAW SEQUENCE LISTING
PATENT APPLICATION *US/09/175,713*

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TIME: 15:35:13

INPUT SET: S29599.raw

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47              (A) LENGTH: 89 amino acids
48              (B) TYPE: amino acid
49              (C) STRANDEDNESS:
50              (D) TOPOLOGY: linear
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52      (ii) MOLECULE TYPE: protein
53
54
55
56
57      (xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:
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62      Cys Leu Ser Asp Gly Lys Pro Val Ser Leu Ser Tyr Arg Cys Pro Cys
63                  20                  25                  30
64
65      Arg Phe Phe Glu Ser His Val Ala Arg Ala Asn Val Lys His Leu Lys
66              35                  40                  45
67
68      Ile Leu Asn Thr Pro Asn Cys Ala Leu Gln Ile Val Ala Arg Leu Lys
69          50                  55                  60
70
71      Asn Asn Asn Arg Gln Val Cys Ile Asp Pro Lys Leu Lys Trp Ile Gln
72      65                  70                  75                  80
73
74      Glu Tyr Leu Glu Lys Ala Leu Asn Lys
75                  85
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82          (C) STRANDEDNESS:
83          (D) TOPOLOGY: linear
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85      (ii) MOLECULE TYPE: protein
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87
88
89
90      (xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:
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94
95      Cys Leu Ser Asp Gly Lys Pro Val Ser Leu Ser Tyr Arg Cys Pro Cys
96                  20                  25                  30
97
98      Arg Phe Phe Glu Ser His Val Ala Arg Ala Asn Val Lys His Leu Lys
99              35                  40                  45
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102            50                  55                  60
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104        Asn Asn Asn Arg Gln Val Cys Ile Asp Pro Lys Leu Lys Trp Ile Gln
105            65                  70                  75                  80
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107        Glu Tyr Leu Glu Lys Ala Leu Asn Lys Arg Phe Lys Met
108                            85                  90
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113            (A) LENGTH: 1856 base pairs
114            (B) TYPE: nucleic acid
115            (C) STRANDEDNESS: double
116            (D) TOPOLOGY: linear
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118    (ii) MOLECULE TYPE: cDNA
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120
121
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123    (xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:
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133    CTGTGCCCTT CAGATTGTAG CCCGGCTGAA GAACAACAAC AGACAAGTGT GCATTGACCC       300
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137    AAAAAGGACT TTCCGCTAGA CCCACTCGAG GAAAACTAAA ACCTTGTGAG AGATGAAAGG       420
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139    GCAAAGACGT GGGGGAGGGG GCCTTAACCA TGAGGACCAG GTGTGTGTGT GGGGTGGGCA       480
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PATENT APPLICATION *US/09/175,713*

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175    TGCGATCCAC AGAAGGTCCT GGTGGTATTT GTAACTTTTT GCAAGGCATT TTTTTATATA     1560
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187    (2) INFORMATION FOR SEQ ID NO:4:
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189         (i) SEQUENCE CHARACTERISTICS:
190              (A) LENGTH: 3526 base pairs
191              (B) TYPE: nucleic acid
192              (C) STRANDEDNESS: double
193              (D) TOPOLOGY: linear
194
195         (ii) MOLECULE TYPE: cDNA
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197
198
199
200         (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:
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202    TCTCCGTCAG CCGCATTGCC CGCTCGGCGT CCGGCCCCCG ACCCGTGCTC GTCCGCCCGC      60
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204    CCGCCCGCCC GCCCGCGCCA TGAACGCCAA GGTCGTGGTC GTGCTGGTCC TCGTGCTGAC     120
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INPUT SET: S29599.raw

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SEQUENCE VERIFICATION REPORT
PATENT APPLICATION   *US/09/175,713*

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